

Bis(2-amino-4-methylpyrimidin-1-i um) hexaaquacobalt(II) disulfate dihydrate

M. Mirzaei,^{a*} H. Eshtiagh-Hosseini,^a S. Zarghami,^a Z. Karrabi,^a M. Saeedi^a and J. T. Mague^{b*}

^aDepartment of Chemistry, Ferdowsi University of Mashhad, 917751436 Mashhad, Iran, and ^bDepartment of Chemistry, Tulane University, New Orleans, LA 70118, USA

Correspondence e-mail: joelt@tulane.edu

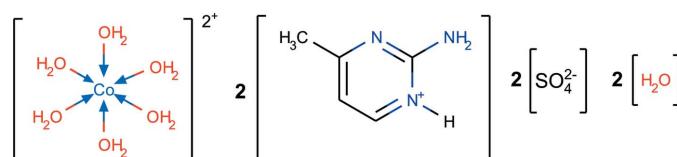
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Key indicators: single-crystal X-ray study; $T = 100\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.002\text{ \AA}$; R factor = 0.026; wR factor = 0.073; data-to-parameter ratio = 19.2.

In the title hydrated mixed-cation salt, $(\text{C}_5\text{H}_8\text{N}_3)_2[\text{Co}(\text{H}_2\text{O})_6](\text{SO}_4)_2 \cdot 2\text{H}_2\text{O}$, the complete octahedral hexaaqua complex cation is generated by crystallographic inversion symmetry. In the crystal, the components are linked by $\text{O}-\text{H}\cdots\text{O}$ and $\text{N}-\text{H}\cdots\text{O}$ hydrogen bonds, the latter, involving pyrimidinium cations and sulfate anions, generating $R_2^2(8)$ loops. These, together with $\pi-\pi$ interactions between centrosymmetrically related pyrimidinium cations [centroid-centroid separation = 3.5460 (8) \AA], lead to the formation of a three-dimensional network.

Related literature

For a report of the structure of the $[\text{Co}(\text{H}_2\text{O})_6]^{2+}$ ion, see: Shiu *et al.* (2004).

**Experimental***Crystal data*

$(\text{C}_5\text{H}_8\text{N}_3)_2[\text{Co}(\text{H}_2\text{O})_6](\text{SO}_4)_2 \cdot 2\text{H}_2\text{O}$

$M_r = 615.49$

Triclinic, $P\bar{1}$

$a = 6.4116 (6)\text{ \AA}$

$b = 7.7751 (7)\text{ \AA}$

$c = 13.0423 (12)\text{ \AA}$

$\alpha = 80.136 (1)^\circ$

$\beta = 80.413 (1)^\circ$

$\gamma = 73.231 (1)^\circ$

$V = 608.57 (10)\text{ \AA}^3$

$Z = 1$

Mo $K\alpha$ radiation

$\mu = 0.96\text{ mm}^{-1}$

$T = 100\text{ K}$

$0.19 \times 0.19 \times 0.12\text{ mm}$

Data collection

Bruker SMART APEX CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 2009)
 $T_{\min} = 0.780$, $T_{\max} = 0.893$

10780 measured reflections
3085 independent reflections
2944 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.025$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.026$
 $wR(F^2) = 0.073$
 $S = 1.09$
3085 reflections

161 parameters
H-atom parameters constrained
 $\Delta\rho_{\max} = 0.59\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.40\text{ e \AA}^{-3}$

Table 1
Selected bond lengths (\AA).

Co1—O1	2.0838 (9)	Co1—O3	2.1140 (10)
Co1—O2	2.0643 (9)		

Table 2
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
O1—H1A \cdots O8	0.84	1.92	2.7562 (14)	171
O1—H1B \cdots O4	0.84	1.97	2.8050 (13)	178
O2—H2A \cdots O4 ⁱ	0.84	1.92	2.7533 (14)	173
O2—H2B \cdots O5	0.84	1.87	2.7077 (13)	174
O3—H3C \cdots O8 ⁱ	0.84	1.92	2.7508 (13)	170
O3—H3D \cdots O7 ⁱⁱ	0.84	1.95	2.7865 (14)	177
N2—H2N \cdots O6 ⁱⁱⁱ	0.91	1.81	2.7155 (15)	172
N3—H3A \cdots O5 ⁱⁱⁱ	0.91	1.87	2.7776 (15)	175
N3—H3B \cdots O6 ^{iv}	0.91	1.98	2.8775 (15)	168
O8—H8A \cdots O7 ⁱⁱ	0.84	1.98	2.7684 (14)	156
O8—H8B \cdots O7 ^v	0.84	2.02	2.8582 (14)	172

Symmetry codes: (i) $x + 1, y, z$; (ii) $x, y + 1, z$; (iii) $-x + 2, -y, -z + 1$; (iv) $-x + 1, -y, -z + 1$; (v) $-x + 1, -y + 1, -z$.

Data collection: *APEX2* (Bruker, 2010); cell refinement: *SAINT* (Bruker, 2009); data reduction: *SAINT*; program(s) used to solve structure: *SHELXM* (Sheldrick, 2008); program(s) used to refine structure: *SHELXTL* (Sheldrick, 2008); molecular graphics: *OLEX2* (Dolomanov *et al.*, 2009); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: HB7029).

References

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Bruker (2010). *APEX2*. Bruker AXS Inc., Madison, Wisconsin, USA.
Dolomanov, O. V., Bourhis, L. J., Gildea, R. J., Howard, J. A. K. & Puschmann, H. (2009). *J. Appl. Cryst.* **42**, 339–341.
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supporting information

Acta Cryst. (2013). E69, m128 [doi:10.1107/S1600536813002146]

Bis(2-amino-4-methylpyrimidin-1-i^{um}) hexaaquacobalt(II) disulfate dihydrate

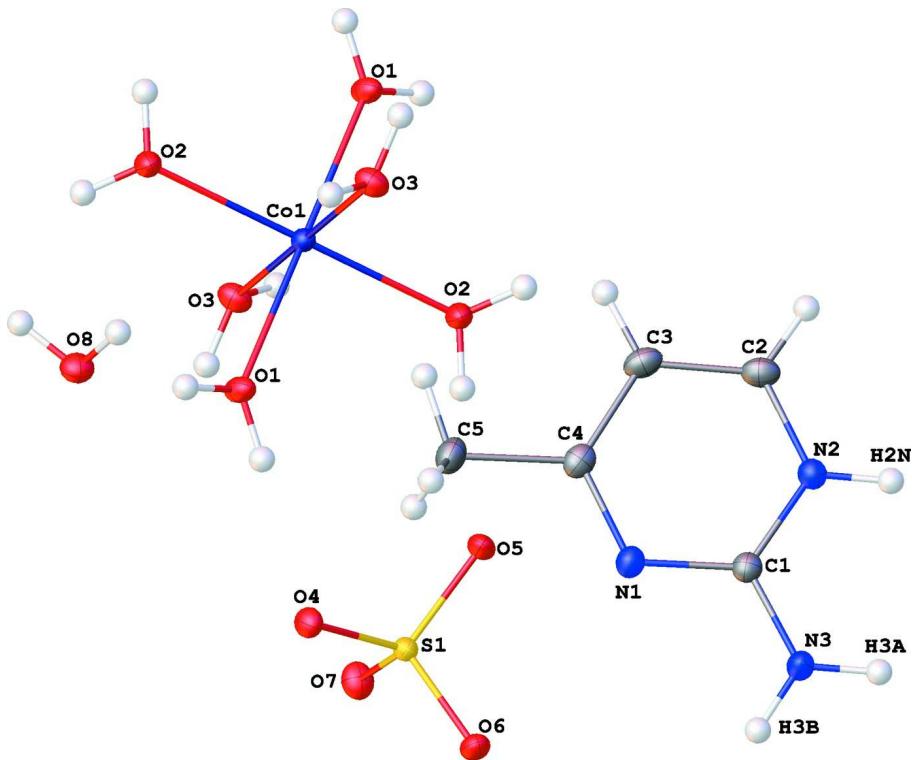
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S1. Comment

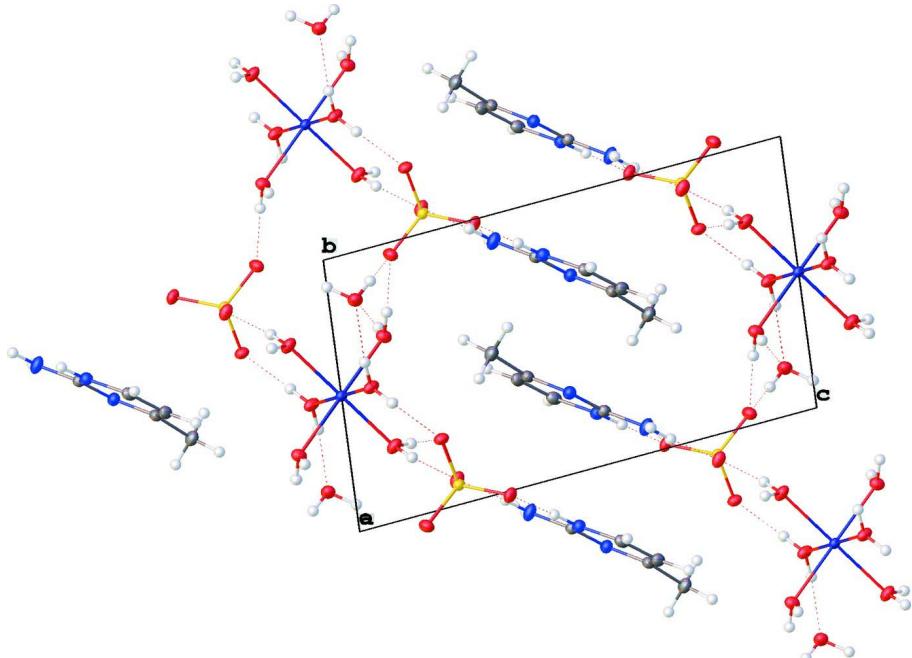
In an attempt to synthesize a cobalt complex of quinoxaline-2,3-dicarboxylic acid by conventional proton transfer processes, cobalt(II) sulfate hexahydrate and the acid were reacted with 2-amino-4-methylpyrimidine. The crystalline product obtained proved to be $[\text{Co}(\text{H}_2\text{O})_6][2\text{a-4m-pym}]_2(\text{SO}_4)_2 \cdot 2\text{H}_2\text{O}$ (2a-4m-pym = 2-amino-4-methylpyrimidinium) with the cobalt cation having crystallographically imposed centrosymmetry (Fig. 1). The Co—O distances range from 2.0643 (9) to 2.1140 (10) Å while the O—Co—O angles range from 90.69 (4) to 93.59 (4)° leading to a somewhat distorted octahedral coordination geometry. The $[\text{Co}(\text{H}_2\text{O})_6]^{2+}$ cations and uncoordinated water molecules form sheets parallel to [110] which are capped on both sides by sulfate ions and held together by O—H···O hydrogen bonds (Fig. 2). N—H···O interactions bind the pyrimidinium cations to the sulfate ions (Fig. 3) with the cations from adjacent sheets intercalating one another (Fig. 2) and stabilized by pairwise $\pi \cdots \pi$ (3.546 Å) interactions between them (center pair of cations in Fig. 2). The N2—H2n···O6 and N3—H3a···O5 hydrogen bonding interactions generate $R_2^2(8)$ synthons.

S2. Experimental

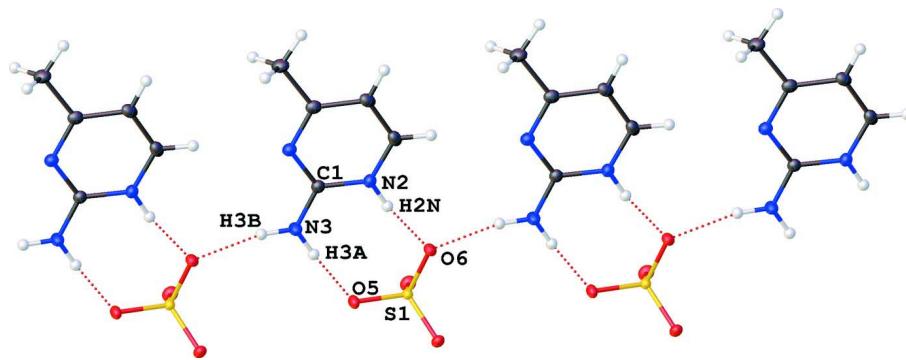
An aqueous solution of cobalt(II) sulfate hexahydrate (0.4 mmol, 0.8 mg) in distilled water (5 ml) was added to an aqueous solution of quinoxaline-2,3-dicarboxylic acid (0.11 mmol, 25 mg) and 2-amino-4-methyl pyrimidine (0.24 mmol, 26 mg). The mixture refluxed for 5 hrs at 75°C. Orange blocks were obtained by slow evaporation of the reaction mixture at room temperature.

**Figure 1**

Perspective view of the title compound with 50% probability ellipsoids.

**Figure 2**

Packing of the title compound viewed down *a*. Hydrogen bonds are indicated by dotted lines.

**Figure 3**

N—H \cdots O hydrogen bonds and the $R_2^2(8)$ synthons (N2, N3, O5 O6).

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Crystal data



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Triclinic, $P\bar{1}$

Hall symbol: -P 1

$a = 6.4116 (6)$ Å

$b = 7.7751 (7)$ Å

$c = 13.0423 (12)$ Å

$\alpha = 80.136 (1)^\circ$

$\beta = 80.413 (1)^\circ$

$\gamma = 73.231 (1)^\circ$

$V = 608.57 (10)$ Å 3

$Z = 1$

$F(000) = 321$

$D_x = 1.679$ Mg m $^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 8977 reflections

$\theta = 2.8\text{--}29.2^\circ$

$\mu = 0.96$ mm $^{-1}$

$T = 100$ K

Block, orange

$0.19 \times 0.19 \times 0.12$ mm

Data collection

Bruker SMART APEX CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

Absorption correction: multi-scan
(SADABS; Sheldrick, 2009)

$T_{\min} = 0.780$, $T_{\max} = 0.893$

10780 measured reflections

3085 independent reflections

2944 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.025$

$\theta_{\max} = 29.2^\circ$, $\theta_{\min} = 2.8^\circ$

$h = -8 \rightarrow 8$

$k = -10 \rightarrow 10$

$l = -17 \rightarrow 17$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.026$

$wR(F^2) = 0.073$

$S = 1.09$

3085 reflections

161 parameters

0 restraints

Primary atom site location: structure-invariant
direct methods

Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites

H-atom parameters constrained

$w = 1/[\sigma^2(F_o^2) + (0.0386P)^2 + 0.2936P]$
where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.001$

$\Delta\rho_{\max} = 0.59$ e Å $^{-3}$

$\Delta\rho_{\min} = -0.40$ e Å $^{-3}$

Special details

Experimental. The diffraction data were obtained from 3 sets of 400 frames, each of width 0.5 °. in omega, collected at phi = 0.00, 90.00 and 180.00 °. and 2 sets of 800 frames, each of width 0.45 ° in phi, collected at omega = -30.00 and 210.00 °. The scan time was 10 sec/frame.

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R-factor wR and goodness of fit S are based on F^2 , conventional R-factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R-factors(gt) etc. and is not relevant to the choice of reflections for refinement. R-factors based on F^2 are statistically about twice as large as those based on F , and R-factors based on ALL data will be even larger. H-atoms attached to carbon were placed in calculated positions (C—H = 0.95 - 0.98 Å) while those attached to nitrogen and oxygen were placed in locations derived from a difference map. All were included as riding contributions with isotropic displacement parameters 1.2 - 1.5 times those of the attached atoms.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Co1	1.0000	0.5000	0.0000	0.00955 (8)
O1	0.68016 (15)	0.50215 (13)	0.06589 (8)	0.01452 (19)
H1A	0.5761	0.5965	0.0592	0.017*
H1B	0.6253	0.4255	0.1061	0.017*
O2	1.13866 (16)	0.27066 (13)	0.09714 (8)	0.0152 (2)
H2A	1.2384	0.2644	0.1331	0.018*
H2B	1.0513	0.2134	0.1317	0.018*
O3	1.01700 (15)	0.67088 (13)	0.10641 (8)	0.01429 (19)
H3C	1.1304	0.7080	0.0967	0.017*
H3D	0.9025	0.7538	0.1193	0.017*
N1	0.77618 (19)	0.29129 (15)	0.48276 (9)	0.0141 (2)
N2	1.13318 (19)	0.19273 (15)	0.53464 (9)	0.0136 (2)
H2N	1.2214	0.1283	0.5837	0.016*
N3	0.8459 (2)	0.09569 (16)	0.63508 (9)	0.0169 (2)
H3A	0.9370	0.0291	0.6825	0.020*
H3B	0.7047	0.0884	0.6443	0.020*
C1	0.9176 (2)	0.19330 (17)	0.55062 (10)	0.0126 (2)
C2	1.2141 (2)	0.28771 (18)	0.44771 (11)	0.0159 (3)
H2	1.3655	0.2840	0.4361	0.019*
C3	1.0775 (2)	0.38846 (18)	0.37710 (11)	0.0164 (3)
H3	1.1307	0.4555	0.3155	0.020*
C4	0.8539 (2)	0.38965 (18)	0.39882 (10)	0.0143 (2)
C5	0.6904 (2)	0.50542 (19)	0.32901 (11)	0.0184 (3)
H5A	0.6319	0.4281	0.2960	0.028*
H5B	0.7619	0.5807	0.2746	0.028*
H5C	0.5701	0.5834	0.3706	0.028*
S1	0.64309 (5)	0.07057 (4)	0.22320 (2)	0.00995 (8)
O4	0.48922 (15)	0.25043 (12)	0.19945 (8)	0.01356 (19)
O5	0.86583 (15)	0.09036 (13)	0.22097 (8)	0.0153 (2)
O6	0.57348 (16)	-0.01584 (13)	0.32903 (8)	0.0148 (2)

O7	0.64425 (16)	-0.04477 (13)	0.14342 (8)	0.0164 (2)
O8	0.36083 (16)	0.82758 (13)	0.05699 (8)	0.01547 (19)
H8A	0.4119	0.8803	0.0939	0.019*
H8B	0.3513	0.8994	0.0009	0.019*

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Co1	0.00874 (12)	0.00915 (12)	0.01048 (13)	-0.00265 (8)	-0.00139 (9)	0.00014 (8)
O1	0.0100 (4)	0.0120 (4)	0.0190 (5)	-0.0025 (3)	-0.0001 (4)	0.0023 (3)
O2	0.0128 (4)	0.0151 (4)	0.0179 (5)	-0.0061 (4)	-0.0050 (4)	0.0048 (4)
O3	0.0107 (4)	0.0149 (4)	0.0178 (5)	-0.0036 (3)	-0.0001 (4)	-0.0047 (4)
N1	0.0149 (5)	0.0136 (5)	0.0133 (5)	-0.0032 (4)	-0.0032 (4)	-0.0005 (4)
N2	0.0125 (5)	0.0137 (5)	0.0144 (5)	-0.0030 (4)	-0.0024 (4)	-0.0013 (4)
N3	0.0140 (5)	0.0206 (6)	0.0157 (6)	-0.0065 (4)	-0.0047 (4)	0.0053 (4)
C1	0.0136 (6)	0.0114 (6)	0.0130 (6)	-0.0029 (4)	-0.0018 (5)	-0.0025 (4)
C2	0.0150 (6)	0.0150 (6)	0.0176 (6)	-0.0052 (5)	0.0031 (5)	-0.0045 (5)
C3	0.0195 (7)	0.0142 (6)	0.0141 (6)	-0.0053 (5)	0.0031 (5)	-0.0022 (5)
C4	0.0187 (6)	0.0119 (6)	0.0119 (6)	-0.0028 (5)	-0.0015 (5)	-0.0033 (4)
C5	0.0222 (7)	0.0171 (6)	0.0155 (6)	-0.0049 (5)	-0.0061 (5)	0.0017 (5)
S1	0.00895 (15)	0.01018 (15)	0.01081 (15)	-0.00313 (11)	-0.00176 (11)	-0.00010 (11)
O4	0.0121 (4)	0.0105 (4)	0.0172 (5)	-0.0015 (3)	-0.0033 (4)	-0.0003 (3)
O5	0.0103 (4)	0.0192 (5)	0.0165 (5)	-0.0067 (4)	-0.0035 (4)	0.0039 (4)
O6	0.0126 (4)	0.0167 (5)	0.0139 (5)	-0.0054 (4)	-0.0010 (4)	0.0033 (4)
O7	0.0178 (5)	0.0149 (5)	0.0171 (5)	-0.0022 (4)	-0.0044 (4)	-0.0054 (4)
O8	0.0165 (5)	0.0142 (4)	0.0172 (5)	-0.0062 (4)	-0.0021 (4)	-0.0024 (4)

Geometric parameters (\AA , ^\circ)

Co1—O1	2.0838 (9)	N3—H3A	0.9099
Co1—O1 ⁱ	2.0838 (9)	N3—H3B	0.9100
Co1—O2	2.0643 (9)	N3—C1	1.3216 (17)
Co1—O2 ⁱ	2.0643 (9)	C2—H2	0.9500
Co1—O3 ⁱ	2.1140 (10)	C2—C3	1.362 (2)
Co1—O3	2.1140 (10)	C3—H3	0.9500
O1—H1A	0.8400	C3—C4	1.412 (2)
O1—H1B	0.8400	C4—C5	1.4935 (19)
O2—H2A	0.8400	C5—H5A	0.9800
O2—H2B	0.8400	C5—H5B	0.9800
O3—H3C	0.8400	C5—H5C	0.9800
O3—H3D	0.8400	S1—O4	1.4772 (10)
N1—C1	1.3503 (17)	S1—O5	1.4749 (9)
N1—C4	1.3352 (17)	S1—O6	1.4851 (10)
N2—H2N	0.9101	S1—O7	1.4834 (10)
N2—C1	1.3621 (17)	O8—H8A	0.8400
N2—C2	1.3558 (17)	O8—H8B	0.8400
O1 ⁱ —Co1—O1	180.0	H3A—N3—H3B	118.7

O1 ⁱ —Co1—O3	89.31 (4)	C1—N3—H3A	121.5
O1—Co1—O3	90.69 (4)	C1—N3—H3B	119.7
O1 ⁱ —Co1—O3 ⁱ	90.69 (4)	N1—C1—N2	121.68 (12)
O1—Co1—O3 ⁱ	89.31 (4)	N3—C1—N1	119.51 (12)
O2 ⁱ —Co1—O1 ⁱ	93.59 (4)	N3—C1—N2	118.82 (12)
O2—Co1—O1	93.59 (4)	N2—C2—H2	120.1
O2 ⁱ —Co1—O1	86.41 (4)	N2—C2—C3	119.73 (13)
O2—Co1—O1 ⁱ	86.41 (4)	C3—C2—H2	120.1
O2—Co1—O2 ⁱ	180.0	C2—C3—H3	121.2
O2 ⁱ —Co1—O3	88.21 (4)	C2—C3—C4	117.61 (12)
O2 ⁱ —Co1—O3 ⁱ	91.79 (4)	C4—C3—H3	121.2
O2—Co1—O3 ⁱ	88.21 (4)	N1—C4—C3	122.45 (12)
O2—Co1—O3	91.79 (4)	N1—C4—C5	116.47 (12)
O3—Co1—O3 ⁱ	180.0	C3—C4—C5	121.07 (12)
Co1—O1—H1A	121.3	C4—C5—H5A	109.5
Co1—O1—H1B	133.1	C4—C5—H5B	109.5
H1A—O1—H1B	105.3	C4—C5—H5C	109.5
Co1—O2—H2A	123.6	H5A—C5—H5B	109.5
Co1—O2—H2B	115.8	H5A—C5—H5C	109.5
H2A—O2—H2B	109.5	H5B—C5—H5C	109.5
Co1—O3—H3C	116.1	O4—S1—O6	110.08 (6)
Co1—O3—H3D	115.8	O4—S1—O7	108.96 (6)
H3C—O3—H3D	112.0	O5—S1—O4	109.75 (6)
C4—N1—C1	117.86 (12)	O5—S1—O6	108.84 (6)
C1—N2—H2N	118.7	O5—S1—O7	109.68 (6)
C2—N2—H2N	120.7	O7—S1—O6	109.52 (6)
C2—N2—C1	120.59 (12)	H8A—O8—H8B	102.1
<hr/>			
N2—C2—C3—C4	0.4 (2)	C2—N2—C1—N3	178.45 (12)
C1—N1—C4—C3	2.59 (19)	C2—C3—C4—N1	-2.6 (2)
C1—N1—C4—C5	-176.26 (12)	C2—C3—C4—C5	176.18 (12)
C1—N2—C2—C3	1.75 (19)	C4—N1—C1—N2	-0.37 (19)
C2—N2—C1—N1	-1.82 (19)	C4—N1—C1—N3	179.37 (12)

Symmetry code: (i) $-x+2, -y+1, -z$.

Hydrogen-bond geometry (\AA , $^{\circ}$)

$D\cdots H\cdots A$	$D\cdots H$	$H\cdots A$	$D\cdots A$	$D\cdots H\cdots A$
O1—H1A \cdots O8	0.84	1.92	2.7562 (14)	171
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N3—H3A \cdots O5 ^{iv}	0.91	1.87	2.7776 (15)	175
N3—H3B \cdots O6 ^v	0.91	1.98	2.8775 (15)	168

O8—H8A…O7 ⁱⁱⁱ	0.84	1.98	2.7684 (14)	156
O8—H8B…O7 ^{vi}	0.84	2.02	2.8582 (14)	172

Symmetry codes: (ii) $x+1, y, z$; (iii) $x, y+1, z$; (iv) $-x+2, -y, -z+1$; (v) $-x+1, -y, -z+1$; (vi) $-x+1, -y+1, -z$.